

**Does Not Comply** 

OIPE

RAW SEQUENCE LISTING DATE: 01/17/2002 PATENT APPLICATION: US/10/032,990 TIME: 15:34:31

Input Set : A:\Seq\_Listing\_-\_P2930R1C11.wpd
Output Set: N:\CRF3\01172002\J032990.raw

```
3 <110> APPLICANT: Botstein, David
                                                                      Corrected Diskette Needed
      4
              Desnoyers, Luc
      5
              Ferrara, Napoleone
      6
              Fong, Sherman
      7
              Gao, Wei-Qiang
      8
              Goddard, Audrey
      9
              Gurney, Austin L.
     10
              Pan, James
     11
              Roy, Margaret Ann
     12
              Stewart, Timothy A.
     13
              Tumas, Daniel
     14
              Watanabe, Colin K.
     15
              Wood, William I.
     17 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
     18
              Acids Encoding the Same
     20 <130> FILE REFERENCE: P2930R1C11
C--> 22 <140> CURRENT APPLICATION NUMBER: US/10/032,990
C--> 22 <141> CURRENT FILING DATE: 2001-12-27
     22 <150> PRIOR APPLICATION NUMBER: 60/095,325
     23 <151> PRIOR FILING DATE: 1998-08-04
     25 <150> PRIOR APPLICATION NUMBER: 60/112,851
     26 <151> PRIOR FILING DATE: 1998-12-16
     28 <150> PRIOR APPLICATION NUMBER: 60/113,145
     29 <151> PRIOR FILING DATE: 1998-12-16
     31 <150> PRIOR APPLICATION NUMBER: 60/113,511
     32 <151> PRIOR FILING DATE: 1998-12-22
     34 <150> PRIOR APPLICATION NUMBER: 60/115,558
     35 <151> PRIOR FILING DATE: 1999-01-12
     37 <150> PRIOR APPLICATION NUMBER: 60/115,565
     38 <151> PRIOR FILING DATE: 1999-01-12
    40 <150> PRIOR APPLICATION NUMBER: 60/115,733
    41 <151> PRIOR FILING DATE: 1999-01-12
    43 <150> PRIOR APPLICATION NUMBER: 60/119,341
    44 <151> PRIOR FILING DATE: 1999-02-09
    46 <150> PRIOR APPLICATION NUMBER: 60/119,537
    47 <151> PRIOR FILING DATE: 1999-02-10
    49 <150> PRIOR APPLICATION NUMBER: 60/119,965
    50 <151> PRIOR FILING DATE: 1999-02-12
    52 <150> PRIOR APPLICATION NUMBER: 60/162,506
    53 <151> PRIOR FILING DATE: 1999-10-29
    55 <150> PRIOR APPLICATION NUMBER: 60/170,262
    56 <151> PRIOR FILING DATE: 1999-12-09
    58 <150> PRIOR APPLICATION NUMBER: 60/187,202
```

Input Set : A:\Seq\_Listing\_-P2930R1C11.wpd
Output Set: N:\CRF3\01172002\J032990.raw

```
59 <151> PRIOR FILING DATE: 2000-03-03
       61 <150> PRIOR APPLICATION NUMBER: PCT/US99/12252
       62 <151> PRIOR FILING DATE: 1999-06-02
       64 <150> PRIOR APPLICATION NUMBER: PCT/US99/28634
       65 <151> PRIOR FILING DATE: 1999-12-01
       67 <150> PRIOR APPLICATION NUMBER: PCT/US99/28551
       68 <151> PRIOR FILING DATE: 1999-12-02
       70 <150> PRIOR APPLICATION NUMBER: PCT/US00/03565
       71 <151> PRIOR FILING DATE: 2000-02-11
       73 <150> PRIOR APPLICATION NUMBER: PCT/US00/04414
       74 <151> PRIOR FILING DATE: 2000-02-22
       76 <150> PRIOR APPLICATION NUMBER: PCT/USQ0/05841
 W--> 77 <151> PRIOR FILING DATE: 2000 (-03-02 delete spoee
       79 <150> PRIOR APPLICATION NUMBER: PCT/US00/08439
       80 <151> PRTOR FILING DATE: 2000-03-30
       82 <150> PRIOR APPLICATION NUMBER: PCT/US00/14941
       83 <151> PRIOR FILING DATE: 2000-05-30
       85 <150> PRIOR APPLICATION NUMBER: PCT/US00/15264
       86 <151> PRIOR FILING DATE: 2000-06-02
       88 <150> PRIOR APPLICATION NUMBER: PCT/US00/32678
       89 <151> PRIOR FILING DATE: 2000-12-01
  W 2 92 (140) CURRENT APPLICATION NUMBER: US 09/866,034
  C--> 93 (141) CURRENT FILING DATE: 2001-05-25
4517 95 <160> NUMBER OF SEQ ID NOS: 38
       97 <210> SEQ ID NO: 1
       98 <211> LENGTH: 1283
       99 <212> TYPE: DNA
       100 <213> ORGANISM: Homo sapiens
       102 <400> SEQUENCE: 1
           eggacgeqtg ggacceatae ttgetggtet gatecatgea caaggegggg 50
           ctgctaggcc tctgtgcccg ggcttggaat tcggtgcgga tggccagctc 100
           cgqqatqacc cgccqggacc cgctcgcaaa taaggtggcc ctggtaacgg 150
       109 cctccaccga cgggatcggc ttcgccatcg cccggcgttt ggcccaggac 200
       111 ggggcccatg tggtcgtcag cagccggaag cagcagaatg tggaccaggc 250
       113 ggtggccacg ctgcaggggg aggggctgag cgtgacgggc accgtgtgcc 300
       115 atgtggggaa ggcggaggac cgggagcggc tggtggccac ggctgtgaag 350
       117 cttcatggag gtatcgatat cctagtctcc aatgctgctg tcaacccttt 400
      119 ctttggaagc ataatggatg tcactgagga ggtgtgggac aagactctgg 450
      121 acattaatgt gaaggcccca gccctgatga caaaggcagt ggtgccagaa 500
      123 atggagaaac gaggaggcgg ctcagtggtg atcgtgtctt ccatagcagc 550
      125 cttcaqtcca tctcctggct tcagtcctta caatgtcagt aaaacagcct 600
       127 tgctgggcct gaccaagacc ctggccatag agetggcccc aaggaacatt 650
      129 aggqtgaact gcctagcacc tggacttatc aagactagct tcagcaggat 700
      131 gctctqqatq qacaaggaaa aagaggaaag catgaaagaa accctgcgga 750
      133 taagaaggtt aggcgagcca gaggattgtg ctggcatcgt gtctttcctg 800
      135 tgctctgaag atgccagcta catcactggg gaaacagtgg tggtgggtgg 850
      137 aggaaccccg tcccgcctct gaggaccggg agacagccca caggccagag 900
      139 ttgggctcta gctcctggtg ctgttcctgc attcacccac tggcctttcc 950
       141 cacctctqct caccttactq ttcacctcat caaatcagtt ctgccctgtg 1000
```

Input Set : A:\Seq\_Listing\_-\_P2930R1C11.wpd
Output Set: N:\CRF3\01172002\J032990.raw

```
143 aaaagatcca gccttccctg ccgtcaaggt ggcgtcttac tcgggattcc 1050
    tgctgttgtt gtggccttgg gtaaaggcct cccctgagaa cacaggacag 1100
     gcctgctgac aaggctgagt ctaccttggc aaagaccaag atatttttc 1150
147
    ctqqqccact gqtqaatctq aqqqqtqatq qqaqaqaaqq aacctqqaqt 1200
149
     ggaaggagca gagttgcaaa ttaacagctt gcaaatgagg tgcaaataaa 1250
    atgcagatga ttgcgcggct ttgaaaaaaa aaa 1283
153
155 <210> SEO ID NO: 2
156 <211> LENGTH: 278
157 <212> TYPE: PRT
158 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 2
     Met His Lys Ala Gly Leu Leu Gly Leu Cys Ala Arg Ala Trp Asn
                                                                15
162
                                           10
164
     Ser Val Arg Met Ala Ser Ser Gly Met Thr Arg Arg Asp Pro Leu
165
                                           25
     Ala Asn Lys Val Ala Leu Val Thr Ala Ser Thr Asp Gly Ile Gly
167
168
                       35
                                           40
170
     Phe Ala Ile Ala Arq Arq Leu Ala Gln Asp Gly Ala His Val Val
171
                       50
                                           55
173
     Val Ser Ser Arg Lys Gln Gln Asn Val Asp Gln Ala Val Ala Thr
174
                       65
                                           70
176
     Leu Gln Gly Glu Gly Leu Ser Val Thr Gly Thr Val Cys His Val
177
                      80
                                           85
179
     Gly Lys Ala Glu Asp Arg Glu Arg Leu Val Ala Thr Ala Val Lys
                                          100
180
                       95
     Leu His Gly Gly Ile Asp Ile Leu Val Ser Asn Ala Ala Val Asn
182
                                          115
183
                      110
     Pro Phe Phe Gly Ser Ile Met Asp Val Thr Glu Glu Val Trp Asp
185
                                          130
186
                      125
188
     Lys Thr Leu Asp : Ile Asn Val Lys Ala Pro Ala Leu Met Thr Lys
                                                               150
189
                      140
     Ala Val Val Pro Glu Met Glu Lys Arg Gly Gly Gly Ser Val Val
191
                                          160
192
                      155
     Ile Val Ser Ser Ile Ala Ala Phe Ser Pro Ser Pro Gly Phe Ser
194
                                          175
                                                               180
195
                      170
     Pro Tyr Asn Val Ser Lys Thr Ala Leu Leu Gly Leu Thr Lys Thr
197
198
                      185
                                          190
     Leu Ala Ile Glu Leu Ala Pro Arg Asn Ile Arg Val Asn Cys Leu
200
                                          205
201
                      200
     Ala Pro Gly Leu Ile Lys Thr Ser Phe Ser Arg Met Leu Trp Met
203
                                          220
                                                               225
204
                      215
     Asp Lys Glu Lys Glu Glu Ser Met Lys Glu Thr Leu Arg Ile Arg
206
207
                      230
                                          235
                                                               240
    Arg Leu Gly Glu Pro Glu Asp Cys Ala Gly Ile Val Ser Phe Leu
209
                                          250
                                                               255
210
                      245
    Cys Ser Glu Asp Ala Ser Tyr Ile Thr Gly Glu Thr Val Val Val
212
                                          265
                                                               270
213
                      260
215
    Gly Gly Gly Thr Pro Ser Arg Leu
216
                      275
```

Input Set : A:\Seq\_\_Listing\_-\_P2930R1C11.wpd
Output Set: N:\CRF3\01172002\J032990.raw

- Output Set: N:\CRF3\01172002\J032990.raw 218 <210> SEQ ID NO: 3 219 <211> LENGTH: 21 220 <212> TYPE: DNA 221 <213> ORGANISM: Artificial Sequence 223 <220> FEATURE: 224 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe 226 <400> SEQUENCE: 3 227 gcataatgga tgtcactgag g 21 229 <210> SEQ ID NO: 4 230 <211> LENGTH: 23 231 <212> TYPE: DNA 232 <213> ORGANISM: Artificial Sequence 234 <220> FEATURE: 235 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe 237 <400> SEQUENCE: 4 238 agaacaatcc tgctgaaagc tag 23 240 <210> SEQ ID NO: 5 241 <211> LENGTH: 46 242 <212> TYPE: DNA 243 <213> ORGANISM: Artificial Sequence 245 <220> FEATURE: 246 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe 248 <400> SEQUENCE: 5 249 gaaacgagga ggcggctcag tggtgatcgt gtcttccata gcagcc 46 251 <210> SEQ ID NO: 6 252 <211> LENGTH: 3121 253 <212> TYPE: DNA 254 <213> ORGANISM: Homo sapiens 256 <400> SEQUENCE: 6 257 gegeeetgag eteegeetee gggeeegata geggeatega gagegeetee 50 259 gtcgaggacc aggcggcgca gggggccggc gggcgaaagg aggatgaggg 100 261 ggcgcagcag ctgctgaccc tgcagaacca ggtggcgcgg ctggaggagg 150 263 agaaccgaga ctttctggct gcgctggagg acgccatgga gcagtacaaa 200 265 ctgcagagcg accggctgcg tgagcagcag gaggagatgg tggaactgcg 250 267 gctgcggtta gagctggtgc ggccaggctg ggggggcctg cggctcctga 300 269 atggcctgcc tcccgggtcc tttgtgcctc gacctcatac agccccctg 350 271 gggggtgecc acgcccatgt gctgggcatg gtgccgcctg cctgcctccc 400 273 tggagatgaa gttggctctg agcagagggg agagcaggtg acaaatggca 450
- gggaggetgg agetgagttg etgaetgagg tgaacagget gggaagtgge 500
  tetteagetg etteagagga ggaagaggag gaggaggage egeecaggeg 550
  gaeettacae etgegeagaa ataggateag eaactgeagt eagagggegg 600
  gggeacgeec agggagtetg eeagagagga agggeecaga getttgeett 650
  gaggagttgg atgeageeat teeagggtee agageagttg gtgggageaa 700
  ggeecgagtt eaggeeegee aggteeeeee tgeeacagee teagagtgge 750
  ggetggeeca ggeecageagaagaegga agagetggetat eaacateege 800
- 289 atgaaggagg agettattgg egagetggte egeacaggaa aggeagetea 850 291 ggeectgaac egecageaca gecagegtat eegggagetg gageaggagg 900 293 cagageaggt gegggeegag etgagtgaag gecagaggea getgegggag 950
- 295 ctcgagggca aggagctcca ggatgctggc gagcggtctc ggctccagga 1000

Input Set : A:\Seq\_\_Listing\_-\_P2930R1C11.wpd
Output Set: N:\CRF3\01172002\J032990.raw

```
gttccgcagg agggtcgctg cggcccagag ccaggtgcag gtgctgaagg 1050
299 aqaaqaaqca qqctacqqaq cqqctqqtqt cactqtcgqc ccaqaqtqag 1100
301 aageqactgc aggagetega geggaaegtg cageteatge ggeageagea 1150
    gggacagctg cagaggcggc ttcgcgagga gacggagcag aagcggcgcc 1200
303
    tqqaqqcaqa aatqaqcaaq cqqcaqcacc qcqtcaaqqa qctqqaqctg 1250
305
307
    aagcatgagc aacagcagaa gatcctgaag attaagacgg aagagatcgc 1300
309
    gqccttccaq aggaagaggc gcagtggcag caacggctct gtggtcagcc 1350
311 tggaacagca gcagaagatt gaggagcaga agaagtggct ggaccaggag 1400
    atggagaagg tgctacagca gcggcgggcg ctggaggagc tggggggagga 1450
313
    gctccacaag cgggaggcca tcctggccaa gaaggaggcc ctgatgcagg 1500
315
    agaagacggg gctggagagc aagcgcctga gatccagcca ggccctcaac 1550
317
319 gaggacateg tgcgagtgtc cagccggctg gagcacctgg agaaggagct 1600
321 qtccqaqaaq aqcqqqcaqc tqcqqcaqqq caqcqcccaq agccaqcaqc 1650
323 agateeqeqq qqaqateqae aqeetqegee aggagaagga etegetgete 1700
325 aagcagegee tggagatega eggeaagetg aggeagggga gtetgetgte 1750
327 ccccqaqqaq qaqcqqacqc tqttccagtt ggatgaggcc atcgaggccc 1800
329 tggatgctgc cattgagtat aagaatgagg ccatcacatg ccgccagcgg 1850
331 qtqcttcqqq cctcaqcctc qttqctqtcc caqtqcqaqa tgaacctcat 1900
    ggccaagete agetacetet cateeteaga gaccagagee etectetgea 1950
335 agtattttga caaggtggtg acgctccgag aggagcagca ccagcagcag 2000
    attgccttct cggaactgga gatgcagctg gaggagcagc agaggctggt 2050
339
    gtactggctg gaggtggccc tggagcggca gcgcctggag atggaccgcc 2100
341 agetgaeeet geageagaag gageaegage agaacatgea getgeteetg 2150
    cagcagagtc gagaccacct cggtgaaggg ttagcagaca gcaggaggca 2200
    gtatgaggee eggatteaag etetggagaa ggaaetggge egttaeatgt 2250
347
    ggataaacca ggaactgaaa cagaagctcg gcggtgtgaa cgctgtaggc 2300
349 cacagcaggg gtggggggaa gaggagcctg tgctcggagg gcagacaggc 2350
    tectggaaat gaagatgage tecacetgge accegagett etetggetgt 2400
351
    cocceteae tgaggggee coccegeacce gggaggagae gegggaettg 2450
353
    gtccacgctc cgttaccctt gacctggaaa cgctcgagcc tgtgtggtga 2500
355
    ggagcagggg tcccccgagg aactgaggca gcgggaggcg gctgagcccc 2550
357
    tggtggggcg ggtgcttcct gtgggtgagg caggcctgcc ctggaacttt 2600
359
    gggcctttgt ccaagccccg gcgggaactg cgacgagcca gcccggggat 2650
363 gattgatgtc cggaaaaacc ccctgtaagc cctcggggca gaccctgcct 2700
365 tggagggaga ctccgagcct gctgaaaggg gcagctgcct gttttgcttc 2750
367 tgtgaagggc agtccttacc gcacacccta aatccaggcc ctcatctgta 2800
369 ccctcactgq qatcaacaaa tttqqqccat gqcccaaaag aactggaccc 2850
371 tcatttaaca aaataatatg caaattccca ccacttactt ccatgaagct 2900
373 gtggtaccca attgccgcct tgtgtcttgc tcgaatctca ggacaattct 2950
375 qqtttcaqqc qtaaatqqat qtqcttqtaq ttcagggqtt tggccaagaa 3000
377
    tcatcacgaa agggtcggtg gcaaccaggt tgtggtttaa atggtcttat 3050
379 gtatataggg gaaactggga gactttagga tcttaaaaaa ccatttaata 3100
381 aaaaaaaatc tttqaaggga c 3121
383 <210> SEO ID NO: 7
384 <211> LENGTH: 830
385 <212> TYPE: PRT
386 <213> ORGANISM: Homo sapiens
388 <400> SEQUENCE: 7
389 Met Glu Gln Tyr Lys Leu Gln Ser Asp Arg Leu Arg Glu Gln Gln
```

VERIFICATION SUMMARY

DATE: 01/17/2002

PATENT APPLICATION: US/10/032,990

TIME: 15:34:32

Input Set : A:\Seq\_Listing\_-\_P2930R1C11.wpd Output Set: N:\CRF3\01172002\J032990.raw

L:22 M:270 C: Current Application Number differs, Replaced Current Application No

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:77 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD L:92 M:280 W: Numeric Identifier already exists, <140> found multiple times

L:92 M:281 W: Numeric Fields not Ordered, <140> not ordered!.

L:92 M:270 C: Current Application Number differs, Replaced Current Application Number

L:93 M:281 W: Numeric Fields not Ordered, <141> not ordered!. L:93 M:271 C: Current Filing Date differs, Replaced Current Filing Date